FIG. 1

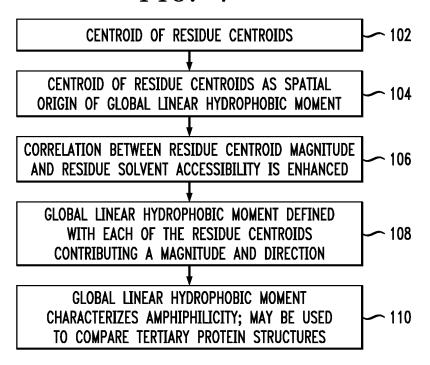
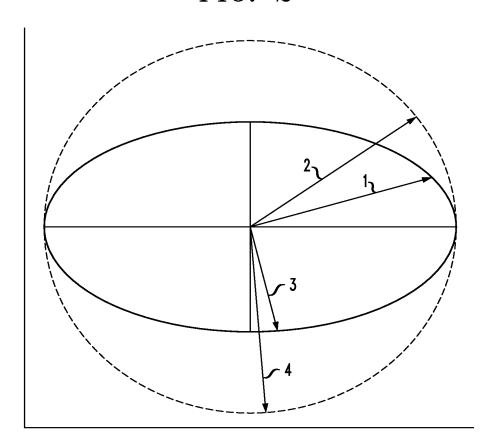


FIG. 2

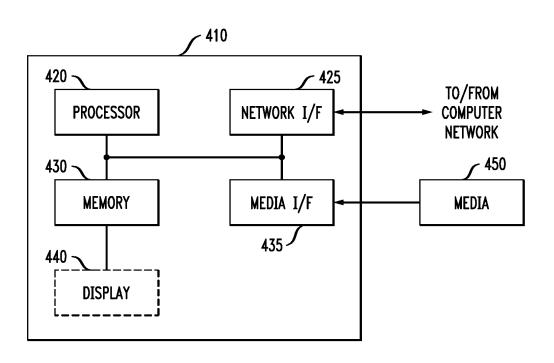


2/7 YOR920030162US1 REPLACEMENT SHEET

CORRELATION COEFFICIENTS OF ELLIPSOIDAL AND RADIAL DISTANCES WITH SOLVENT ACCESSIBILITIES

93	1.39	<u> </u>	. 69.	1.59	1.63	04.	1.31	1.31	2.55	1.48	1.25	1.78	1.69	2.41	8	1.52	1.39	1.61	1.45	1.38	1.31	1.73	5.09
92	1.03		1.28	1.33	1.39	=	1:21	1.22	2.48	1.40	1.17	1.63	1.60	2.23	1.48	1.19	==	1.21	1.28	1.07	1.14	1.63	1.88 8.
RADIAL	0.644	0.685	0.686	0.677	0.703	0.712	0.735	0.719	0.455	0.534	0.726	0.623	0.641	0.473	0.481	0.612	0.620	0.524	0.674	0.535	0.667	0.559	0.487
ELLIPSOIDAL	0.672 0.632	0.756	0.734	0.736	0.760	0.744	0.754	0.733	0.647	0.620	0.739	0.690	0.715	0.625	0.578	0.655	0.651	0.573	0.703	0.568	0.685	0.640	0.565
RESIDUE Number	208 213	216 218	219	223	228	245	256	257	271	536	300	309	318	338	348	351	356	405	448	449	468	200	574
PROTEIN	1AUN 1LBU	1YAL 2ACT	1EUG	1AKZ	三	1TH	166V	1BN	2DRI	1AUA	1A3H	1LDM	1FSZ	1KFU	1UBY	1A26	1CIN	1PHC	1GAI	186V	3PBG	3C0X	田
93	1.66	1.50 2.04	1.55	2.60	2.89	2.21	1.99	1.86	1.45	1.51	1.48	1.79	1.61	1.26	1.56	1.35	1.66	1.20	1.83	1.43	1.47	2.03	1.36
92	1.56	1.31	6.	2.40	2.48	2.16	1.87	1.75	1.25	1.34	1.18	1.50	1.19	1.14	1.13	1.12	1.56	1.13	1.44	1.13	1.23	2.00	1.18
RADIAL	0.795	0.776 0.669	0.700	0.575	0.517	0.653	0.686	0.682	0.800	0.757	0.737	0.608	9/9'0	0.825	0.670	0.795	0.762	0.742	0.666	0.632	0.722	0.669	0.783
ELLIPSOIDAL	0.852	0.821	0.769	0.752	0.702	0.815	0.783	0.792	0.831	0.788	0.780	0.689	0.728	0.847	0.724	0.818	0.817	0.763	0.704	0.670	0.759	0.765	0.805
RESIDUE NUMBER	96 105	108	116	116	120	123	123	129	122	141	145	147	151	151	153	154	160	166	170	186	195	206	506
PROTEIN	1CDZ 2RAC	1 1 1 1 1 1	1DLW	1BL0	1DZ0	10TS	1A4V	3121	1PD0	2SNS	1AT0	1H97	1A6M	1118	1002	1PHR	1CZT	121P	1E6C	16KY	2KFZ	1DZV	되

FIG. 4



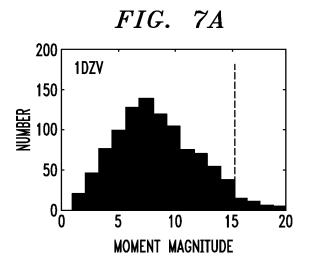
4/7 YOR920030162US1 REPLACEMENT SHEET

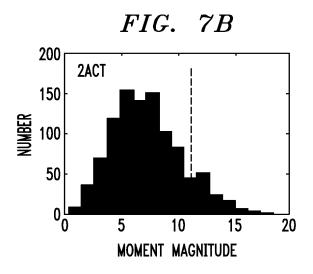
FIG.	5		MOMENT	MAGN	MEAN HYDROPHOBICITIES	PHOBICITIES			
PROTEIN	MOMENT MAGNITUDE	RANDOM MAGNITUDE	NUMBER GREATER	MEAN HYDROPHOBICITY	PROTEIN	MOMENT MAGNITUDE	RANDOM MAGNITUDE	NUMBER Greater	MEAN HYDROPHOBICIT
1007	6.16	9.49	765	-0.083	1AUN	3.57	6.54	898	-0.122
2RAC	4.24	7.91	864	-0.090	1LBU	3.49	6.82	878	-0.135
1,00	7.62	8.71	280	-0.222	1YAL	6.58	7.32	565	-0.158
IN IN	8.91	10.24	575	-0.083	2ACT	11.20	7.29	112	-0.092
1DLW	5.97	7.68	229	-0.044	1506	5.51	7.41	695	-0.064
1BL0	5.13	10.18	869	-0.164	1AKZ	1.58	7.59	989	-0.105
1020	6.29	9.80	762	-0.224	10DH	6.34	7.39	605	-0.065
10TS	3.13	10.66	977	-0.063	昰	6.92	7.00	491	-0.109
1A4V	9.95	10.64	533	-0.089	166V	6.93	7.71	260	-0.154
3121	10.35	8.43	269	-0.139	1BN	6.38	7.62	623	-0.158
1 5 6 1	8.87	8.40	414	-0.047	2DRI	3.26	8.86	943	-0.145
2SNS	6.31	8.15	671	-0.227	1AUA	17.09	7.86	ယ	-0.134
1AT0	10.11	8.02	242	-0.069	1A3H	1.83	6.57	975	-0.131
1197	5.22	8.94	831	-0.110	MQ1	13.81	7.90	જ	-0.039
1A6W	3.41	8.73	943	-0.117	1FSZ	4.83	7.48	784	-0.089
1118	3.92	8.76	915	-0.150	1KFU	11.05	8.88	281	-0.118
1002	3.79	8.47	922	-0.122	10BY	8.84	8.27	412	-0.104
罢	5.20	8.04	770	-0.121	1A26	5.44	7.44	725	-0.117
1021	9.91	9.16	395	-0.120	1CTN	2.78	6.73	937	-0.124
121P	7.68	7.80	478	-0.134	문	3.67	7.23	88	-0.087
1E6C	9.54	8.71	366	-0.054	1GAI	5.79	6.07	522	-0.114
16KY	9.63	8.43	332	-0.170	186	7.75	6.40	536	-0.100
2KFZ	6.82	8.07	594	960.0-	3PBG	5.82	6.34	244	-0.121
1DZV	15.31	8.41	40	-0.042	3COX	4.32	6.95	908	-0.120
10CH	10.39	7.90	232	-0.094	臣	6.92	09.7	267	-0.129

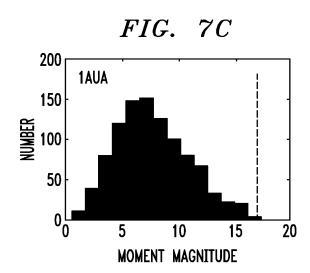
FIG. 5

FIG. 6
NEUMAIER HYDROPHOBICITY SCALE

AMINO ACID	HYDROPHOBICITY	AMINO ACID	HYDROPHOBICITY
LYS	-1.00	HIS	-0.15
ASP	-0.97	ALA	-0.06
GLU	-0.85	TYR	0.35
ARG	-0.80	CYS	0.56
GLN	-0.71	TRP	0.57
ASN	-0.70	MET	0.68
SER	-0.48	VAL	0.75
PRO	-0.45	LEU	0.83
THR	-0.38	PHE	0.99
GLY	-0.32	ILE	1.00







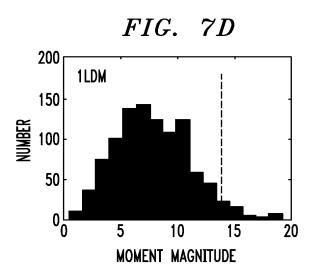


FIG. 8

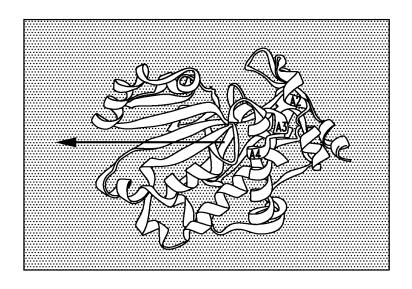
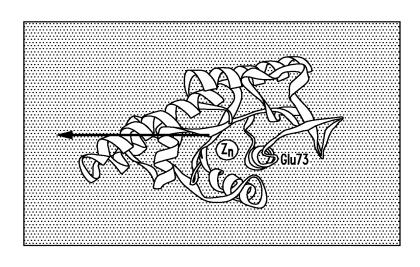


FIG. 9



7/7 YOR920030162US1 REPLACEMENT SHEET

 $FIG.\ 10$ defensin and defensin like moment magnitudes and mean hydrophobicities

PROTEIN	MOMENT MAGNITUDE	RANDOM MAGNITUDE	NUMBER Greater	MEAN HYDOPHOBICITY	g ₂	gź
1AHL	16.86	9.09	24	-0.089	1.85	2.02
1APF	23.16	10.55	31	-0.100	1.20	2.66
1B8W	15.79	18.98	539	-0.182	1.97	5.34
1BNB	13.58	17.00	615	-0.001	3.13	4.96
1DFN	3.34	9.25	955	0.070	2.29	2.44
1FD3	5.74	9.35	806	-0.079	1.97	2.36
1FJN	7.02	12.37	745	-0.076	1.64	3.96
1SH1	36.35	13.60	9	-0.184	1.98	3.60

FIG. 11A

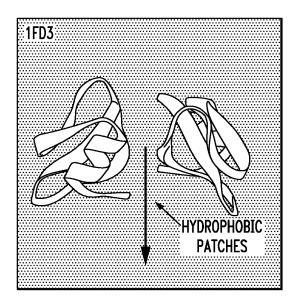


FIG. 11B

